

09806311 .071801



PCT09

**Does Not Comply
Corrected Diskette Needed**

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/806,311B

DATE: 07/16/2002

TIME: 15:39:26

Input Set : A:\PTO.VSK.txt

Output Set: N:\CRF3\07162002\I806311B.raw

3 <110> APPLICANT: Macina, Roberto A.
 5 <120> TITLE OF INVENTION: A NOVEL METHOD OF DIAGNOSING, MONITORING, STAGING, IMAGING
 AND TREATING
 6 GASTROINTESTINAL CANCERS
 8 <130> FILE REFERENCE: DEX-0184
 C--> 10 <140> CURRENT APPLICATION NUMBER: US/09/806,311B
 C--> 10 <141> CURRENT FILING DATE: 2002-04-30
 10 <150> PRIOR APPLICATION NUMBER: PCT/US99/22725
 11 <151> PRIOR FILING DATE: 1999-09-30
 13 <150> PRIOR APPLICATION NUMBER: 60/102,879
 14 <151> PRIOR FILING DATE: 1998-10-02
 16 <160> NUMBER OF SEQ ID NOS: 2
 18 <170> SOFTWARE: PatentIn version 3.1

ERRORED SEQUENCES

93 <210> SEQ ID NO: 2
 94 <211> LENGTH: 153
 95 <212> TYPE: PRT
 96 <213> ORGANISM: Homo sapien
 98 <400> SEQUENCE: 2
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 101 1 5 10 15
 104 Gly Asp Ile Ile Met Arg Pro Ser Cys Ala Pro Gly Trp Phe Tyr His
 105 20 25 30
 108 Lys Ser Asn Cys Tyr Gly Tyr Phe Arg Lys Leu Arg Asn Trp Ser Asp
 109 35 40 45
 112 Ala Glu Leu Glu Cys Gln Ser Tyr Gly Asn Gly Ala His Leu Ala Ser
 113 50 55 60
 116 Ile Leu Ser Leu Lys Glu Ala Ser Thr Ile Ala Glu Tyr Ile Ser Gly
 117 65 70 75 80
 120 Tyr Gln Arg Ser Gln Pro Ile Trp Ile Gly Leu His Asp Pro Gln Lys
 121 85 90 95
 124 Arg Gln Gln Trp Gln Trp Ile Asp Gly Ala Met Tyr Leu Tyr Arg Ser
 125 100 105 110
 128 Trp Ser Gly Lys Ser Met Gly Gly Asn Lys His Cys Ala Glu Met Ser
 129 115 120 125
 132 Ser Asn Asn Asn Phe Leu Thr Trp Ser Ser Asn Glu Cys Asn Lys Arg
 133 130 135 140
 136 Gln His Phe Leu Cys Lys Tyr Arg Pro
 137 145 150
 E--> 143 (3) ~Delete
 E--> 149 (3)

09806311 - 0716001

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/806,311B

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Input Set : A:\PTO.VSK.txt

Output Set: N:\CRF3\07162002\I806311B.raw

L:10 M:270 C: Current Application Number differs, Replaced Current Application No

L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:90 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:1200

L:143 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:2

M:332 Repeated in SeqNo=2